



## SEQUENCE LISTING

&lt;110&gt; Seiki Motoharu

&lt;120&gt; DNA CODING FOR NOVEL POLIPEPTIDE

&lt;130&gt; 1241.18

&lt;140&gt; 09/806,232

&lt;141&gt; 2001-03-28

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&lt;150&gt; JP10-276258

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&lt;150&gt; JP10-291505

&lt;151&gt; 1998-09-29

&lt;160&gt; 28

&lt;170&gt; PatentIn Ver. 2.0

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&lt;212&gt; PRT

&lt;213&gt; Mouse

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cgc	ccc	gtc	tcc	gac	ttc	agc	ctc	ccg	cct	ggc	ggc	atc	gac	gct	gcc	1410
Arg	Pro	Val	Ser	Asp	Phe	Ser	Leu	Pro	Pro	Gly	Gly	Ile	Asp	Ala	Ala	
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Phe	Ser	Trp	Ala	His	Asn	Asp	Arg	Thr	Tyr	Phe	Phe	Lys	Asp	Gln	Leu	
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Tyr	Trp	Arg	Tyr	Asp	Asp	His	Thr	Arg	His	Met	Asp	Pro	Gly	Tyr	Pro	
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 Trp Thr Ala Ala Gln Ala Leu Thr Leu  
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Pro	Gly	Tyr	Pro	Lys 485	Pro	Ile	Thr	Val	Trp 490	Lys	Gly	Ile	Pro	Gln 495	Ala					
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 Tyr Gly Tyr Leu Leu Pro Tyr Glu Ser Arg Ala Ser Ala Leu His Ser  
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Ile	Thr	Tyr	Ser	Ile	His	Asn	Tyr	Thr	Pro	Lys	Val	Gly	Glu	Leu	Asp		
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 Pro Arg Cys Gly Val Pro Asp Lys Phe Gly Ala Glu Ile Lys Ala Asn Val Arg  
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 Arg Lys Arg Tyr Ala Ile Gln Gly Leu Lys Trp Gln His Asn Glu Ile Thr Phe  
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 Cys Ile Gln Asn Tyr Thr Pro Lys Val Gly Glu Tyr Ala Thr Tyr Glu Ala Ile  
 130 135 140  
 Arg Lys Ala Phe Arg Val Trp Glu Ser Ala Thr Pro Leu Arg Phe Arg Glu Val  
 145 150 155 160  
 Pro Tyr Ala Tyr Ile Arg Glu Gly His Glu Lys Gln Ala Asp Ile Met Ile Phe  
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 Phe Ala Glu Gly Phe His Gly Asp Ser Thr Pro Phe Asp Gly Glu Gly Gly Phe  
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 Leu Ala His Ala Tyr Phe Pro Gly Pro Asn Ile Gly Gly Asp Thr His Phe Asp  
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 Ser Ala Glu Pro Trp Thr Val Arg Asn Glu Asp Leu Asn Gly Asn Asp Ile Phe  
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 Leu Val Ala Val His Glu Leu Gly His Ala Leu Gly Leu Glu His Ser Ser Asp  
 235 240 245 250  
 Pro Ser Ala Ile Met Ala Pro Phe Tyr Gln Trp Met Asp Thr Glu Asn Phe Val  
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	Trp	Arg	Gly	Leu	Pro	Ala	Ser	Ile	Asn	Thr	Ala	Tyr	Glu	Arg	Lys	Asp	Gly	Lys
	Phe	Val	Phe	Phe	Lys	Gly	Asp	Lys	His	Trp	Val	Phe	Asp	Glu	Ala	Ser	Leu	Glu
380	Pro	Gly	Tyr	Pro	Lys	His	Ile	Lys	Glu	Leu	Gly	Arg	Gly	Leu	Pro	Thr	Asp	Lys
	Ile	Asp	Ala	Ala	Leu	Phe	Trp	Met	Pro	Asn	Gly	Lys	Thr	Tyr	Phe	Phe	Arg	Gly
415	Asn	Lys	Tyr	Tyr	Arg	Phe	Asn	Glu	Glu	Leu	Arg	Ala	Val	Asp	Ser	Glu	Tyr	Pro
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470	Asn	Asn	Gln	Lys	Leu	Lys	Val	Glu	Pro	Gly	Tyr	Pro	Lys	Ser	Ala	Leu	Arg	Asp
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40	Asn	Trp	Leu	Arg	Leu	Tyr	Gly	Tyr	Leu	Pro	Gln	Pro	Ser	Arg	His	Met	Ser	Thr
55	Met	Arg	Ser	Ala	Gln	Ile	Leu	Ala	Ser	Ala	Leu	Ala	Glu	Met	Gln	Arg	Phe	Tyr
75	Gly	Ile	Pro	Val	Thr	Gly	Val	Leu	Asp	Glu	Glu	Thr	Lys	Glu	Trp	Met	Lys	Arg
95	Pro	Arg	Cys	Gly	Val	Pro	Asp	Gln	Phe	Gly	Val	Arg	Val	Lys	Ala	Asn	Leu	Arg
110	Arg	Arg	Arg	Lys	Arg	Tyr	Ala	Leu	Thr	Gly	Arg	Lys	Trp	Asn	Asn	His	His	Leu
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145	Ala	Val	Arg	Arg	Ala	Phe	Arg	Val	Trp	Glu	Gln	Ala	Thr	Pro	Leu	Val	Phe	Gln
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Lys	Pro	Glu	Arg	Pro	Pro	Lys	Pro	Gly	Pro	Pro	Val	Gln	Pro	Arg	Ala	Thr	Glu
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Arg	Pro	Asp	Gln	Tyr	Gly	Pro	Asn	Ile	Cys	Asp	Gly	Asp	Phe	Asp	Thr	Val	Ala
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Met	Leu	Arg	Gly	Glu	Met	Phe	Val	Phe	Lys	Gly	Arg	Trp	Phe	Trp	Arg	Val	Arg
	380					385					390					395	
His	Asn	Arg	Val	Leu	Asp	Asn	Tyr	Pro	Met	Pro	Ile	Gly	His	Phe	Trp	Arg	Gly
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Pro	Gln	Pro	Leu	Thr	Ser	Tyr	Gly	Leu	Gly	Ile	Pro	Tyr	Asp	Arg	Ile	Asp	Thr
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Trp	Arg	Phe	Asn	Glu	Glu	Thr	Gln	Arg	Gly	Asp	Pro	Gly	Tyr	Pro	Lys	Pro	Ile
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	505				510					515					520		
Ala	Ala	Tyr	Thr	Tyr	Phe	Tyr	Lys	Gly	Thr	Lys	Tyr	Trp	Lys	Phe	Asp	Asn	Glu
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Arg	Leu	Arg	Met	Glu	Pro	Gly	Tyr	Pro	Lys	Ser	Ile	Leu	Arg	Asp	Phe	Met	Gly
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		615					620					625					630
Val	Pro	Leu	Leu	Leu	Leu	Leu	Cys	Val	Leu	Gly	Leu	Thr	Tyr	Ala	Leu	Val	Gln
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 <213> Homo sapiens

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Thr	Glu	Gln	Tyr	Phe	Asn	Val	Glu	Val	Trp	Leu	Gln	Lys	Tyr	Gly	Tyr	Leu	Pro
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Pro	Thr	Asp	Pro	Arg	Met	Ser	Val	Leu	Arg	Ser	Ala	Glu	Thr	Met	Gln	Ser	Ala
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Asn	Thr	Ile	Asp	Trp	Met	Lys	Lys	Pro	Arg	Cys	Gly	Val	Pro	Asp	Gln	Thr	Arg
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145					150					155					160		
Thr	Pro	Leu	Thr	Phe	Glu	Glu	Val	Pro	Tyr	Ser	Glu	Leu	Glu	Asn	Gly	Lys	Arg
		165					170					175					180
Asp	Val	Asp	Ile	Thr	Ile	Ile	Phe	Ala	Ser	Gly	Phe	His	Gly	Asp	Ser	Ser	Pro
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Lys	Asp	Gln	Trp	Phe	Trp	Arg	Val	Arg	Asn	Asn	Arg	Val	Met	Asp	Gly	Tyr	Pro
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Thr	Tyr	Phe	Phe	Lys	Gly	Asp	Arg	Tyr	Trp	Arg	Tyr	Ser	Glu	Glu	Met	Lys	Thr
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Glu	Gly	His	Ser	Pro	Pro	Asp	Asp	Val	Asp	Ile	Val	Ile	Lys	Leu	Asp	Asn	Thr
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Ala	Ser	Thr	Val	Lys	Ala	Ile	Ala	Ile	Val	Ile	Pro	Cys	Ile	Leu	Ala	Leu	Cys
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Leu	Leu	Val	Leu	Val	Tyr	Thr	Val	Phe	Gln	Phe	Lys	Arg	Lys	Gly	Thr	Pro	Arg
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 <213> Homo sapiens

<400> 26

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Gly	Val	Glu	Trp	Leu	Ser	Arg	Phe	Gly	Tyr	Leu	Pro	Pro	Ala	Asp	Pro	Thr	Thr
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Phe	Gly	Gly	Leu	Glu	Ala	Thr	Gly	Ile	Leu	Asp	Glu	Ala	Thr	Leu	Ala	Leu	Met
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Arg	Arg	Gln	Ala	Pro	Ala	Pro	Thr	Lys	Trp	Asn	Lys	Arg	Asn	Leu	Ser	Trp	Arg
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Val	Arg	Thr	Phe	Pro	Arg	Asp	Ser	Pro	Leu	Gly	His	Asp	Thr	Val	Arg	Ala	Leu
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Met	Tyr	Tyr	Ala	Leu	Lys	Val	Trp	Ser	Asp	Ile	Ala	Pro	Leu	Asn	Phe	His	Glu
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Val	Ala	Gly	Ser	Thr	Ala	Asp	Ile	Gln	Ile	Asp	Phe	Ser	Lys	Ala	Asp	His	Asn
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His	His	His	Thr	Ala	Gly	Tyr	Thr	His	Phe	Asn	Asp	Asp	Glu	Ala	Trp	Thr	Phe
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Arg	Ser	Ser	Asp	Ala	His	Gly	Met	Asp	Leu	Phe	Ala	Val	Ala	Val	His	Glu	Phe
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Tyr	Tyr	Gln	Gly	Pro	Val	Gly	Asp	Pro	Leu	Arg	Tyr	Gly	Leu	Pro	Tyr	Glu	Asp
				275					280					285			
Lys	Val	Arg	Val	Trp	Gln	Leu	Tyr	Gly	Val	Arg	Glu	Ser	Val	Ser	Pro	Thr	Ala
290					295						300				305		
Gln	Pro	Glu	Glu	Pro	Pro	Leu	Leu	Pro	Glu	Pro	Pro	Asp	Asn	Arg	Ser	Ser	Ala
			310					315					320				
Pro	Pro	Arg	Lys	Asp	Val	Pro	His	Arg	Cys	Ser	Thr	His	Phe	Asp	Ala	Val	Ala
325					330				335						340		
Gln	Ile	Arg	Gly	Glu	Ala	Phe	Phe	Phe	Lys	Gly	Lys	Tyr	Phe	Trp	Arg	Leu	Thr
		345					350					355				360	
Arg	Asp	Arg	His	Leu	Val	Ser	Leu	Gln	Pro	Ala	Gln	Met	His	Arg	Phe	Trp	Arg
				365				370						375			
Gly	Leu	Pro	Leu	His	Leu	Asp	Ser	Val	Asp	Ala	Val	Tyr	Glu	Arg	Thr	Ser	Asp
380					385						390					395	
His	Lys	Ile	Val	Phe	Phe	Lys	Gly	Asp	Arg	Tyr	Trp	Val	Phe	Lys	Asp	Asn	Asn
			400					405					410				
Val	Glu	Glu	Gly	Tyr	Pro	Arg	Pro	Val	Ser	Asp	Phe	Ser	Leu	Pro	Pro	Gly	Gly
415					420					425					430		
Ile	Asp	Ala	Ala	Phe	Ser	Trp	Ala	His	Asn	Asp	Arg	Thr	Tyr	Phe	Phe	Lys	Asp
		435					440					445				450	
Gln	Leu	Tyr	Trp	Arg	Tyr	Asp	Asp	His	Thr	Arg	His	Met	Asp	Pro	Gly	Tyr	Pro
				455				460						465			
Ala	Gln	Ser	Pro	Leu	Trp	Arg	Gly	Val	Pro	Ser	Thr	Leu	Asp	Asp	Ala	Met	Arg
470					475						480					485	
Trp	Ser	Asp	Gly	Ala	Ser	Tyr	Phe	Phe	Arg	Gly	Gln	Glu	Tyr	Trp	Lys	Val	Leu
			490					495					500				
Asp	Gly	Glu	Leu	Glu	Val	Ala	Pro	Gly	Tyr	Pro	Gln	Ser	Thr	Ala	Arg	Asp	Trp
505					510					515					520		
Leu	Val	Cys	Gly	Asp	Ser	Gln	Ala	Asp	Gly	Ser	Val	Ala	Ala	Gly	Val	Asp	Ala

Ala	Glu	Gly	Pro	Arg	Ala	Pro	Pro	Gly	Gln	His	Asp	Gln	Ser	Arg	Ser	Glu	Asp
				545					550					555			
Gly	Tyr	Glu	Val	Cys	Ser	Cys	Thr	Ser	Gly	Ala	Ser	Ser	Pro	Pro	Gly	Ala	Pro
	560					565					570					575	
Gly	Pro	Leu	Val	Ala	Ala	Thr	Met	Leu	Leu	Leu	Leu	Pro	Pro	Leu	Ser	Pro	Gly
			580					585					590				
Ala	Leu	Trp	Thr	Ala	Ala	Gln	Ala	Leu	Thr	Leu							
595					600					605							

<210> 27  
 <211> 645  
 <212> PRT  
 <213> Homo sapiens

Met	Pro	Arg	Ser	Arg	Gly	Gly	Arg	Ala	Ala	Pro	Gly	Pro	Pro	Pro	Pro	Pro	Pro
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	20					25					30					35	
Leu	Leu	Leu	Leu	Pro	Ala	Leu	Cys	Cys	Leu	Pro	Gly	Ala	Ala	Arg	Ala	Ala	Ala
			40					45					50				
Ala	Ala	Ala	Gly	Ala	Gly	Asn	Arg	Ala	Ala	Val	Ala	Val	Ala	Val	Ala	Arg	Ala
	55				60					65					70		
Asp	Glu	Ala	Glu	Ala	Pro	Phe	Ala	Gly	Gln	Asn	Trp	Leu	Lys	Ser	Tyr	Gly	Tyr
	75					80					85					90	
Leu	Leu	Pro	Tyr	Asp	Ser	Arg	Ala	Ser	Ala	Leu	His	Ser	Ala	Lys	Ala	Leu	Gln
				95					100					105			
Ser	Ala	Val	Ser	Thr	Met	Gln	Gln	Phe	Tyr	Gly	Ile	Pro	Val	Thr	Gly	Val	Leu
	110				115						120					125	
Asp	Gln	Thr	Thr	Ile	Glu	Trp	Met	Lys	Lys	Pro	Arg	Cys	Gly	Val	Pro	Asp	His
			130					135					140				
Pro	His	Leu	Ser	Arg	Arg	Arg	Asn	Lys	Arg	Tyr	Ala	Leu	Thr	Gly	Gln	Lys	
	145				150				155					160			
Trp	Arg	Gln	Lys	His	Ile	Thr	Tyr	Ser	Ile	His	Asn	Tyr	Thr	Pro	Lys	Val	Gly
		165				170						175					180
Glu	Leu	Asp	Thr	Arg	Lys	Ala	Ile	Arg	Gln	Ala	Phe	Asp	Val	Trp	Gln	Lys	Val
				185				190					195				
Thr	Pro	Leu	Thr	Phe	Glu	Glu	Val	Pro	Tyr	His	Glu	Ile	Lys	Ser	Asp	Arg	Lys
	200				205						210					215	
Glu	Ala	Asp	Ile	Met	Ile	Phe	Phe	Ala	Ser	Gly	Phe	His	Gly	Asp	Ser	Ser	Pro
			220					225					230				
Phe	Asp	Gly	Glu	Gly	Gly	Phe	Leu	Ala	His	Ala	Tyr	Phe	Pro	Gly	Pro	Gly	Ile
	235				240				245					250			
Gly	Gly	Asp	Thr	His	Phe	Asp	Ser	Asp	Glu	Pro	Trp	Thr	Leu	Gly	Asn	Ala	Asn
		255				260						265					270
His	Asp	Gly	Asn	Asp	Leu	Phe	Leu	Val	Ala	Val	His	Glu	Leu	Gly	His	Ala	Leu
				275					280					285			
Gly	Leu	Glu	His	Ser	Ser	Asp	Pro	Ser	Ala	Ile	Met	Ala	Pro	Phe	Tyr	Gln	Tyr
	290					295					300					305	
Met	Glu	Thr	His	Asn	Phe	Lys	Leu	Pro	Gln	Asp	Asp	Leu	Gln	Gly	Ile	Gln	Lys
			310						315				320				
Ile	Tyr	Gly	Pro	Pro	Ala	Glu	Pro	Leu	Glu	Pro	Thr	Arg	Pro	Leu	Pro	Thr	Leu
					330					335					340		
Pro	Val	Arg	Arg	Ile	His	Ser	Pro	Ser	Glu	Arg	Lys	His	Glu	Arg	Gln	Pro	Arg
		345					350					355					360
Pro	Pro	Arg	Pro	Pro	Leu	Gly	Asp	Arg	Pro	Ser	Thr	Pro	Gly	Thr	Lys	Pro	Asn
				365					370					375			
Ile	Cys	Asp	Gly	Asn	Phe	Asn	Thr	Val	Ala	Leu	Phe	Arg	Gly	Glu	Met	Phe	Val
	380					385					390					395	
Phe	Lys	Asp	Arg	Trp	Phe	Trp	Arg	Leu	Arg	Asn	Asn	Arg	Val	Gln	Glu	Gly	Tyr
			400					405					410				
Pro	Met	Gln	Ile	Glu	Gln	Phe	Trp	Lys	Gly	Leu	Pro	Ala	Arg	Ile	Asp	Ala	Ala
	415				420					425					430		



Tyr	Glu	Arg	Ala	Asp	Gly	Arg	Phe	Val	Phe	Phe	Lys	Gly	Asp	Lys	Tyr	Trp	Val
		435					440					445					450
Phe	Lys	Glu	Val	Thr	Val	Glu	Pro	Gly	Tyr	Pro	His	Ser	Leu	Gly	Glu	Leu	Gly
				455					460					465			
Ser	Cys	Leu	Pro	Arg	Glu	Gly	Ile	Asp	Thr	Ala	Leu	Arg	Trp	Glu	Pro	Val	Gly
	470					475					480					485	
Lys	Thr	Tyr	Phe	Phe	Lys	Gly	Glu	Arg	Tyr	Trp	Arg	Tyr	Ser	Glu	Glu	Arg	Arg
			490					495					500				
Ala	Thr	Asp	Pro	Gly	Tyr	Pro	Lys	Pro	Ile	Thr	Val	Trp	Lys	Gly	Ile	Pro	Gln
505					510					515					520		
Ala	Pro	Gln	Gly	Ala	Phe	Ile	Ser	Lys	Glu	Gly	Tyr	Tyr	Thr	Tyr	Phe	Tyr	Lys
		525					530					535					540
Gly	Arg	Asp	Tyr	Trp	Lys	Phe	Asp	Asn	Gln	Lys	Leu	Ser	Val	Glu	Pro	Gly	Tyr
				545					550					555			
Pro	Arg	Asn	Ile	Leu	Arg	Asp	Trp	Met	Gly	Cys	Asn	Gln	Lys	Glu	Val	Glu	Arg
	560					565					570					575	
Arg	Lys	Glu	Arg	Arg	Leu	Pro	Gln	Asp	Asp	Val	Asp	Ile	Met	Val	Thr	Ile	Asn
			580					585					590				
Asp	Val	Pro	Gly	Ser	Val	Asn	Ala	Val	Ala	Val	Val	Ile	Pro	Cys	Ile	Leu	Ser
595					600				605						610		
Leu	Cys	Ile	Leu	Val	Leu	Val	Tyr	Thr	Ile	Phe	Gln	Phe	Lys	Asn	Lys	Thr	Gly
		615					620					625					630
Pro	Gln	Pro	Val	Thr	Tyr	Tyr	Lys	Arg	Pro	Val	Gln	Glu	Trp	Val			
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<210> 28  
 <211> 618  
 <212> PRT  
 <213> Mouse

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Gly	Trp	Arg	Ala	Pro	Gly	Arg	Leu	Leu	Pro	Leu	Leu	Pro	Ala	Leu	Cys	Cys	Leu
	20					25					30					35	
Ala	Ala	Ala	Ala	Gly	Ala	Gly	Lys	Pro	Ala	Gly	Ala	Asp	Ala	Pro	Phe	Ala	Gly
			40					45					50				
Gln	Asn	Trp	Leu	Lys	Ser	Tyr	Gly	Tyr	Leu	Leu	Pro	Tyr	Glu	Ser	Arg	Ala	Ser
	55				60					65					70		
Ala	Leu	His	Ser	Gly	Lys	Ala	Leu	Gln	Ser	Ala	Val	Ser	Thr	Met	Gln	Gln	Phe
		75					80					85					90
Tyr	Gly	Ile	Pro	Val	Thr	Gly	Val	Leu	Asp	Gln	Thr	Thr	Ile	Glu	Trp	Met	Lys
				95					100					105			
Lys	Pro	Arg	Cys	Gly	Val	Pro	Asp	His	Pro	His	Leu	Ser	Arg	Arg	Arg	Arg	Asn
	110					115					120				125		
Lys	Arg	Tyr	Ala	Leu	Thr	Gly	Gln	Lys	Trp	Arg	Gln	Lys	His	Ile	Thr	Tyr	Ser
			130					135					140				
Ile	His	Asn	Tyr	Thr	Pro	Lys	Val	Gly	Glu	Leu	Asp	Thr	Arg	Lys	Ala	Ile	Arg
	145				150					155					160		
Gln	Ala	Phe	Asp	Val	Trp	Gln	Lys	Val	Thr	Pro	Leu	Thr	Phe	Glu	Glu	Val	Pro
		165				170					175					180	
Tyr	His	Glu	Ile	Lys	Ser	Asp	Arg	Lys	Glu	Ala	Asp	Ile	Met	Ile	Phe	Phe	Ala
				185					190					195			
Ser	Gly	Phe	His	Gly	Asp	Ser	Ser	Pro	Phe	Asp	Gly	Glu	Gly	Gly	Phe	Leu	Ala
	200					205					210					215	
His	Ala	Tyr	Phe	Pro	Gly	Pro	Gly	Ile	Gly	Gly	Asp	Thr	His	Phe	Asp	Ser	Asp
			220					225					230				
Glu	Pro	Trp	Thr	Leu	Gly	Asn	Ala	Asn	His	Asp	Gly	Asn	Asp	Leu	Phe	Leu	Val
	235				240					245					250		
Ala	Val	His	Glu	Leu	Gly	His	Ala	Leu	Gly	Leu	Glu	His	Ser	Asn	Asp	Pro	Ser
		255					260					265					270
Ala	Ile	Met	Ala	Pro	Phe	Tyr	Gln	Tyr	Met	Glu	Thr	His	Asn	Phe	Lys	Leu	Pro
				275					280					285			
Gln	Asp	Asp	Leu	Gln	Gly	Ile	Gln	Lys	Ile	Tyr	Gly	Pro	Pro	Ala	Glu	Pro	Leu

290						295				300					305
Glu	Pro	Thr	Arg	Pro	Leu	His	Thr	Leu	Pro	Val	Arg	Arg	Ile	His	Ser
			310					315					320		
Glu	Arg	Lys	His	Glu	Arg	His	Pro	Arg	Pro	Pro	Arg	Pro	Pro	Leu	Gly
325					330					335					340
Pro	Ser	Thr	Pro	Gly	Ala	Lys	Pro	Asn	Ile	Cys	Asp	Gly	Asn	Phe	Asn
		345					350				355				360
Ala	Leu	Phe	Arg	Gly	Glu	Met	Phe	Val	Phe	Lys	Asp	Arg	Trp	Phe	Trp
				365				370						375	
Arg	Asn	Asn	Arg	Val	Gln	Glu	Gly	Tyr	Pro	Met	Gln	Ile	Glu	Gln	Phe
	380				385						390				395
Gly	Leu	Pro	Ala	Arg	Ile	Asp	Ala	Ala	Tyr	Glu	Arg	Ala	Asp	Gly	Arg
			400				405					410			
Phe	Phe	Lys	Gly	Asp	Lys	Tyr	Trp	Val	Phe	Lys	Glu	Val	Thr	Val	Glu
415					420					425					430
Tyr	Pro	His	Ser	Leu	Gly	Glu	Leu	Gly	Ser	Cys	Leu	Pro	Arg	Glu	Gly
		435					440					445			450
Thr	Ala	Leu	Arg	Trp	Glu	Pro	Val	Gly	Lys	Thr	Tyr	Phe	Phe	Lys	Gly
				455				460						465	
Tyr	Trp	Cys	Tyr	Ser	Glu	Glu	Arg	Arg	Ala	Thr	Asp	Pro	Gly	Tyr	Pro
	470					475					480				485
Ile	Thr	Val	Trp	Lys	Gly	Ile	Pro	Gln	Ala	Pro	Gln	Gly	Ala	Phe	Ile
			490					495					500		
Glu	Gly	Tyr	Tyr	Thr	Tyr	Phe	Tyr	Lys	Gly	Arg	Asp	Tyr	Trp	Lys	Phe
505					510					515					520
Gln	Lys	Leu	Ser	Val	Glu	Pro	Gly	Tyr	Pro	Arg	Asn	Ile	Leu	Arg	Asp
		525					530					535			540
Gly	Cys	Lys	Gln	Lys	Glu	Val	Glu	Arg	Arg	Lys	Glu	Arg	Arg	Leu	Pro
				545				550						555	
Asp	Val	Asp	Ile	Met	Val	Thr	Ile	Asp	Asp	Val	Pro	Gly	Ser	Val	Asn
	560					565				570					575
Ala	Val	Val	Val	Pro	Cys	Thr	Leu	Ser	Leu	Cys	Leu	Leu	Val	Leu	Leu
			580				585					590			
Ile	Phe	Gln	Phe	Lys	Asn	Lys	Ala	Gly	Pro	Gln	Pro	Val	Thr	Tyr	Tyr
595					600					605					610
Pro	Val	Gln	Glu	Trp	Val										
		615													